SEQUENCE LISTING



110> Michael F. Thomashow Sarah George Fowler Jonathan Vogel Daniel Zarka

<120> Transcription Factors to Improve Plant Stress Tolerance
<130> 21835-00004
<140> US/10/632,436
<141> 2003-08-01
<150> 60/400,777
<151> 2002-08-02
<160> 2
<210> 1
<211> 1308
<212> DNA
<213> Arabidopsis thaliana
<220>
<221><221><221><222></223> This is a cDNA sequence derived from mRNA encoding the putative

<223> This is a cDNA sequence derived from mRNA encoding the putative AP2 domain transcription factor RAV1 (locus tag: Atlg13260). The RAV1 coding sequence consists of nucleotides 92 through 1126 as numbered below.

<400> 1

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gtatacatat acacaacata attcacaaca caacacaaac acatttctgt tttctccatt
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gtttcaaacc ataaaaaaa acacagatta aatggaatcg agtagcgttg atgagagtac
                                                                      120
tacaagtaca ggttccatct gtgaaacccc ggcgataact ccggcgaaaa agtcgtcggt
                                                                      180
aggtaactta tacaggatgg gaagcggatc aagcgttgtg ttagattcag agaacggcgt
                                                                      240
                                                                      300
agaagctgaa tctaggaagc ttccgtcgtc aaaatacaaa ggtgtggtgc cacaaccaaa
cggaagatgg ggagctcaga tttacgagaa acaccagcgc gtgtggctcg ggacattcaa
                                                                      360
cgaagaagac gaagccgctc gtgcctacga cgtcgcggtt cacaggttcc gtcgccgtga
                                                                      420
cgccgtcaca aatttcaaag acgtgaagat ggacgaagac gaggtcgatt tcttgaattc
                                                                      480
tcattcgaaa tctgagatcg ttgatatgtt gaggaaacat acttataacg aagagttaga
                                                                      540
gcagagtaaa cggcgtcgta atggtaacgg aaacatgact aggacgttgt taacgtcggg
                                                                      600
gttgagtaat gatggtgttt ctacgacggg gtttagatcg gcggaggcac tgtttgagaa
                                                                      660
ageggtaacg ccaagegacg ttgggaaget aaacegtttg gttatacega aacatcaege
                                                                      720
                                                                      780
agagaaacat tttccgttac cgtcaagtaa cgtttccgtg aaaggagtgt tgttgaactt
tgaggacgtt aacgggaaag tgtggaggtt ccgttactcg tattggaaca gtagtcagag
                                                                      840
ttatgttttg actaaaggtt ggagcaggtt cgttaaggag aagaatctac gtgctggtga
                                                                      900
cgtggttagt ttcagtagat ctaacggtca ggatcaacag ttgtacattg ggtggaagtc
                                                                      960
gagatccggg tcagatttag atgcgggtcg ggttttgaga ttgttcggag ttaacatttc
                                                                      1020
accggagagt tcaaqaaacg acgtcgtagg aaacaaaaga gtgaacgata ctgagatgtt
                                                                      1080
atcqttqqtq tqtaqcaaqa aqcaacqcat ctttcacqcc tcqtaacaac tcttcttctt
                                                                      1140
tttttttttt tttqttqttt taataatttt taaaaactcc attttcqttt tctttatttq
                                                                      1200
categgtttc tttcttcttg tttaccaaag gttcatgagt tgtttttgtt gtattgatga
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actgtaaatt ttatttatag gataaatttt aaaaagggtt acttagat
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<210> 2 <211> 816

<212> DNA

<213> Arabidopsis thaliana

<220>

<221>

<222>

<223> This is a cDNA sequence derived from mRNA encoding the putative zinc- finger protein Zat12 (locus tag: At5g59820). The Zat12 coding sequence consists of nucleotides 102 through 590 as numbered below.

<400> 2

| atcatcacaa | ctactatcac | accaaactca | aaaaacacaa | accacaagag | gatcatttca | 60 |
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| ttttttattg | tttcgtttta | atcatcatca | tcagaagaaa | aatggttgcg | atatcggaga | 120 |
| | | acggcggcga | | | | 180 |
| aagaaaacgt | tgacggtggc | gatcaaaaac | gcgttttcac | atgtaaaacg | tgtttgaagc | 240 |
| agtttcattc | gttccaagcc | ttaggaggtc | accgtgcgag | tcacaagaag | cctaacaacg | 300 |
| acgctttgtc | gtctggattg | atgaagaagg | tgaaaacgtc | gtcgcatcct | tgtcccatat | 360 |
| gtggagtgga | gtttccgatg | ggacaagctt | tgggaggaca | catgaggaga | cacaggaacg | 420 |
| agagtggggc | tgctggtggc | gcgttggtta | cacgcgcttt | gttgccggag | cccacggtga | 480 |
| ctacgttgaa | gaaatctagc | agtgggaaga | gagtggcttg | tttggatctg | agtctaggga | 540 |
| tggtggacaa | tttgaatctc | aagttggagc | ttggaagaac | agtttattga | ttttatttat | 600 |
| tttccttaaa | ttttctgaat | atatttgttt | ctctcattct | ttgaattttt | cttaatattc | 660 |
| tagattatac | atacatccgc | agatttagga | aactttcata | gagtgtaatc | ttttctttct | 720 |
| gtaaaaatat | attttacttg | tagcattgga | gatttgttat | gagattatct | tacttagcat | 780 |
| ttagtgaata | atctattagc | ctattttgcc | gacgtg | | | 816 |